

## SEQUENCE LISTING

<110> David, WAGNER H

<120> METHODS FOR PREDICTING DEVELOPMENT OF AUTO-IMMUNE DISEASES AND  
TREATMENT OF SAME

<130> 059742-5001-WO

<150> US 60/484,655

<151> 2003-07-07

<150> PCT/US2004/021646

<151> 2004-07-07

<160> 21

<170> PatentIn version 3.3

<210> 1

<211> 6545

<212> DNA

<213> Homo sapiens

<400> 1

|  |     |
|--|-----|
| gagagcagag aacacacttt gccttctctt tggtattgag taatatcaac caaattgcag  | 60  |
| acatctcaac actttggcca ggcagcctgc tgagcaaggt acctcagcca gcatggcagc  | 120 |
| ctctttccca cccaccttgg gactcagttc tgccccagat gaaattcagc acccacatat  | 180 |
| taaattttca gaatggaaat ttaagctgtt ccgggtgaga tcctttgaaa agacacctga  | 240 |
| agaagctcaa aaggaaaaga aggattcctt tgaggggaaa ccctctctgg agcaatctcc  | 300 |
| agcagtcctg gacaaggctg atggtcagaa gccagtccca actcagccat tgttaaaagc  | 360 |
| ccaccctaag ttttcaaaga aatttcacga caacgagaaa gcaagaggca aagcgatcca  | 420 |
| tcaagccaac cttcgacatc tctgccgcat ctgtgggaat tcttttagag ctgatgagca  | 480 |
| caacaggaga tatccagtcc atggtcctgt ggatggtaaa accctaggcc ttttacgaaa  | 540 |
| gaaggaaaag agagctactt cctggccgga cctcattgcc aaggttttcc ggatcgatgt  | 600 |
| gaaggcagat gttgactcga tccacccac tgagttctgc cataactgct ggagcatcat   | 660 |
| gcacaggaag ttttagcagtg ccccatgtga ggtttacttc ccgaggaacg tgaccatgga | 720 |
| gtggcacccc cacacacat cctgtgacat ctgcaacact gcccgtcggg gactcaagag   | 780 |
| gaagagtctt cagccaaact tgcagctcag caaaaaactc aaaactgtgc ttgaccaagc  | 840 |
| aagacaagcc cgtcagcgca agagaagagc tcaggcaagg atcagcagca aggatgtcat  | 900 |
| gaagaagatc gccaaactga gtaagataca tcttagtacc aagctccttg cagtggactt  | 960 |

|            |            |            |             |             |             |      |
|------------|------------|------------|-------------|-------------|-------------|------|
| cccagagcac | tttgtgaaat | ccatctcctg | ccagatctgt  | gaacacattc  | tggtgaccc   | 1020 |
| tgtggagacc | aactgtaagc | atgtcttttg | ccgggtctgc  | attctcagat  | gcctcaaagt  | 1080 |
| catgggcagc | tattgtccct | cttgccgata | tccatgcttc  | cctactgacc  | tggagagtcc  | 1140 |
| agtgaagtcc | tttctgagcg | tcttgaattc | cctgatggtg  | aaatgtccag  | caaaagagtg  | 1200 |
| caatgaggag | gtcagtttgg | aaaaatataa | tcaccacatc  | tcaagtcaca  | aggaatcaaa  | 1260 |
| agagatTTTT | gtgcacatta | ataaaggggg | ccggccccgc  | caacatcttc  | tgctgctgac  | 1320 |
| tccgagagct | cagaagcacc | ggctgagggg | gctcaagctg  | caagtcaaag  | cctttgctga  | 1380 |
| caaagaagaa | ggtggagatg | tgaagtccgt | gtgcatgacc  | ttgttcctgc  | tggtctctgag | 1440 |
| ggcgaggaat | gagcacaggc | aagctgatga | gctggaggcc  | atcatgcagg  | gaaagggctc  | 1500 |
| tggcctgcag | ccagctgttt | gcttggccat | ccgtgtcaac  | accttcctca  | gctgcagtca  | 1560 |
| gtaccacaag | atgtacagga | ctgtgaaagc | catcacaggg  | agacagattt  | ttcagccttt  | 1620 |
| gcatgccctt | cggaatgctg | agaaggtact | tctgccaggc  | taccaccact  | ttgagtggca  | 1680 |
| gccacctctg | aagaatgtgt | cttcagcac  | tgatgttggc  | attattgatg  | ggctgtctgg  | 1740 |
| actatcatcc | tctgtggatg | attaccaggt | ggacaccatt  | gcaaagaggt  | tccgctatga  | 1800 |
| ttcagctttg | gtgtctgctt | tgatggacat | ggaagaagac  | atcttggaag  | gcatgagatc  | 1860 |
| ccaagacctt | gatgattacc | tgaatggccc | cttcaactgtg | gtgggtgaagg | agtcttgtga  | 1920 |
| tggaatggga | gacgtgagtg | agaagcatgg | gagtgggcct  | gtagttccag  | aaaaggcagt  | 1980 |
| ccgtttttca | ttcacaatca | tgaaaattac | tattgcccac  | agctctcaga  | atgtgaaagt  | 2040 |
| atTTgaagaa | gccaaaccta | actctgaact | gtgttgcaag  | ccattgtgcc  | ttatgctggc  | 2100 |
| agatgagtct | gaccacgaga | cgctgactgc | catcctgagt  | cctctcattg  | ctgagagggg  | 2160 |
| ggccatgaag | agcagtgaat | taatgcttga | gctgggaggc  | attctccgga  | ctttcaagtt  | 2220 |
| catcttcagg | ggcaccggct | atgatgaaaa | acttgtgcgg  | gaagtggaag  | gcctcgaggc  | 2280 |
| ttctggctca | gtctacattt | gtactctttg | tgatgccacc  | cgtctggaag  | cctctcaaaa  | 2340 |
| tcttgtcttc | cactctataa | ccagaagcca | tgctgagaac  | ctggaacggt  | atgaggtctg  | 2400 |
| gcgttccaac | ccttaccatg | agtctgtgga | agaactgcgg  | gatcgggtga  | aaggggtctc  | 2460 |
| agctaaacct | ttcattgaga | cagtcccttc | catagatgca  | ctccactgtg  | acattggcaa  | 2520 |
| tgcagctgag | ttctacaaga | tcttccagct | agagataggg  | gaagtgtata  | agaatcccaa  | 2580 |
| tgcttccaaa | gaggaaagga | aaaggtggca | ggccacactg  | gacaagcatc  | tccggaagaa  | 2640 |

|   |      |
|---|------|
| gatgaacctc aaaccaatca tgaggatgaa tggcaacttt gccaggaagc tcatgaccaa | 2700 |
| agagactgtg gatgcagttt gtgagttaat tccttccgag gagaggcacg aggctctgag | 2760 |
| ggagctgatg gatctttacc tgaagatgaa accagtatgg cgatcatcat gccctgctaa | 2820 |
| agagtgccca gaatccctct gccagtacag tttcaattca cagcgttttg ctgagctcct | 2880 |
| ttctacgaag ttcaagtata ggtatgaggg aaaaatcacc aattattttc acaaaacctt | 2940 |
| ggcccatgtt cctgaaatta ttgagagggg tggctccatt ggggcatggg caagtgaggg | 3000 |
| aaatgagtct ggtaacaaac tgtttagggc cttccggaaa atgaatgccg ggcagtccaa | 3060 |
| atgctatgag atggaagatg tcctgaaaca ccactgggtg tacacctcca aatacctcca | 3120 |
| gaagtttatg aatgctcata atgcattaaa aacctctggg tttaccatga accctcaggc | 3180 |
| aagcttaggg gacccattag gcatagagga ctctctggaa agccaagatt caatggaatt | 3240 |
| ttaagtaggg caaccactta tgagttgggt tttgcaattg agtttccttc tgggttgcat | 3300 |
| tgagggtctc tcctagcacc ctttactgct gtgtatgggg cttcaccatc caagaggtgg | 3360 |
| taggttgag taagatgcta cagatgctct caagtcagga atagaaactg atgagctgat  | 3420 |
| tgcttgaggc ttttagtgag ttccgaaaag caacaggaaa aatcagttat ctgaaagctc | 3480 |
| agtaactcag aacaggagta actgcagggg accagagatg agcaaagatc tgtgtgtgtt | 3540 |
| ggggagctgt catgtaaatc aaagccaagg ttgtcaaaga acagccagtg aggccagaaa | 3600 |
| ttggtcttgt ggttttcatt tttttcccc ttgattgatt atattttgta ttgagatatg  | 3660 |
| ataagtgcct tctatttcat ttttgaataa ttcttcattt ttataatttt acatatcttg | 3720 |
| gcttgctata taagattcaa aagagctttt taaatttttc taataatatc ttacatttgt | 3780 |
| acagcatgat gacctttaca aagtgctctc aatgcattta cccattcgtt atataaatat | 3840 |
| gttacatcag gacaactttg agaaaatcag tcctttttta tgtttaaatt atgtatctat | 3900 |
| tgtaaccttc agagtttagg aggtcatctg ctgtcatgga tttttcaata atgaatttag | 3960 |
| aatacacctg ttagctacag ttagttatta aatcttctga taatatatgt ttacttagct | 4020 |
| atcagaagcc aagtatgatt ctttatTTTT actttttcat ttcaagaaat ttagagtttc | 4080 |
| caaatttaga gcttctgcat acagtcttaa agccacagag gcttgtaaaa atataggtta | 4140 |
| gcttgatgtc taaaaatata tttcatgtct tactgaaaca ttttgccaga ctttctccaa | 4200 |
| atgaaacctg aatcaatttt tctaaatcta ggtttcatag agtcctctcc tctgcaatgt | 4260 |
| gttattcttt ctataatgat cagtttactt tcagtggatt cagaattgtg tagcaggata | 4320 |

|            |            |            |             |             |            |      |
|------------|------------|------------|-------------|-------------|------------|------|
| accttgatt  | tttccatccg | ctaagtttag | atggagtcca  | aacgcagtac  | agcagaagag | 4380 |
| ttaacattta | cacagtgcct | tttaccactg | tggaatgttt  | tcacactcat  | ttttccttac | 4440 |
| aacaattctg | aggagtaggt | gttggtatta | tctccatttg  | atgggggttt  | aatgatttgc | 4500 |
| tcaaagtcac | ttaggggtaa | taaatacttg | gcttggaat   | ttaacacagt  | ccttttgtct | 4560 |
| ccaaagccct | tcttctttcc | accacaaatt | aatcactatg  | tttataagggt | agtatcagaa | 4620 |
| tttttttagg | attcacaact | aatcactata | gcacatgacc  | ttgggattac  | atttttatgg | 4680 |
| ggcaggggta | agcggctttt | aaatcatttg | tgtgctctgg  | ctcttttgat  | agaagaaagc | 4740 |
| aacacaaaag | ctccaaagg  | cccctaacc  | ctcttggtgg  | tccagttatt  | tggaactat  | 4800 |
| gatctgcac  | cttaggaatc | tggtatttgc | cagttgctgg  | caatgtagag  | caggcatgga | 4860 |
| attttatatg | ctagtgcac  | ataatgat   | gttagtggt   | attagttttt  | cttcctttga | 4920 |
| ttttatttgc | cataattgct | actcttcata | cacagtatat  | caaagagctt  | gataatttag | 4980 |
| ttgtcaaaag | tgcacggcg  | acattatctt | taattgtatg  | tatttggtgc  | ttcttcaggg | 5040 |
| attgaactca | gtatctttca | ttaaaaaaca | cagcagtttt  | ccttgctttt  | tatatgcaga | 5100 |
| atatcaaagt | catttcta   | ttagttgtca | aaaacatata  | catattttta  | cattagtttt | 5160 |
| tttgaaaact | cttggttttg | tttttttgga | aatgagtggg  | ccactaagcc  | acactttccc | 5220 |
| ttcatcctgc | ttaatccttc | cagcatgtct | ctgcactaat  | aaacagctaa  | attcacataa | 5280 |
| tcaccttatt | tactgaagca | tggtcatgct | ggtttataga  | ttttttaccc  | atttctactc | 5340 |
| tttttctcta | ttggtggcac | tgtaaatact | ttccagtatt  | aaattatcct  | tttctaacac | 5400 |
| tgtaggaact | attttgaatg | catgtgacta | agagcatgat  | ttatagcaca  | acctttccaa | 5460 |
| taatccctta | atcagatcac | attttgataa | accctgggaa  | catctggctg  | caggaatttc | 5520 |
| aatatgtaga | aacgctgcct | atgggttttt | gcccttactg  | ttgagactgc  | aatatcctag | 5580 |
| accctagttt | tatactagag | ttttattttt | agcaatgcct  | attgcaagtg  | caattatata | 5640 |
| ctccagggaa | attcaccaca | ctgaatcgag | catttggtgtg | tgtatgtgtg  | aagtatatct | 5700 |
| gggacttcag | aagtgcaatg | tatttttctc | ctgtgaaacc  | tgaatctaca  | agttttctgc | 5760 |
| caagccactc | aggtgcattg | cagggaccag | tgataatggc  | tgatgaaaat  | tgatgattgg | 5820 |
| tcagtgcagt | caaaaggagc | cttgggatta | ataaacatgc  | actgagaagc  | aagaggagga | 5880 |
| gaaaaagatg | tctttttctt | ccaggtgaac | tggaatttag  | ttttgcctca  | gatttttttc | 5940 |
| ccacaagata | cagaagaaga | taaagatttt | tttggttgag  | agtgtgggtc  | ttgcattaca | 6000 |

```

tcaaacagag ttcaaattcc acacagataa gaggcaggat atataagcgc cagtggtagt 6060
tgaggaggaat aaaccattat ttggatgcag gtgggttttg attgcaaata tgtgtgtgtc 6120
ttcagtgatt gtatgacaga tgatgtattc ttttgatgtt aaaagatttt aagtaagagt 6180
agatacattg tacccatttt acattttctt attttaacta cagtaatcta cataaatata 6240
cctcagaaat cttttttggt gattattttt tgtttttag aattgcactt cagtttattt 6300
tcttacaat aaccttacat tttgtttaat ggcttccaag agcctttttt tttttgtatt 6360
tcagagaaaa ttcaggtacc aggatgcaat ggatttattt gattcagggg acctgtattt 6420
ccatgtcaaa tgttttcaaa taaaatgaaa tatgagtttc aatacttttt atattttaat 6480
atttccttaa tattatgggt attgtccgcc attttgttgt atattgtaaa taaagtttag 6540
attgt 6545

```

```

<210> 2
<211> 1043
<212> PRT
<213> Homo sapiens

```

<400> 2

```

Met Ala Ala Ser Phe Pro Pro Thr Leu Gly Leu Ser Ser Ala Pro Asp
1           5           10           15

```

```

Glu Ile Gln His Pro His Ile Lys Phe Ser Glu Trp Lys Phe Lys Leu
          20           25           30

```

```

Phe Arg Val Arg Ser Phe Glu Lys Thr Pro Glu Glu Ala Gln Lys Glu
          35           40           45

```

```

Lys Lys Asp Ser Phe Glu Gly Lys Pro Ser Leu Glu Gln Ser Pro Ala
          50           55           60

```

```

Val Leu Asp Lys Ala Asp Gly Gln Lys Pro Val Pro Thr Gln Pro Leu
65           70           75           80

```

```

Leu Lys Ala His Pro Lys Phe Ser Lys Lys Phe His Asp Asn Glu Lys
          85           90           95

```

```

Ala Arg Gly Lys Ala Ile His Gln Ala Asn Leu Arg His Leu Cys Arg
          100          105          110

```

Ile Cys Gly Asn Ser Phe Arg Ala Asp Glu His Asn Arg Arg Tyr Pro  
115 120 125

Val His Gly Pro Val Asp Gly Lys Thr Leu Gly Leu Leu Arg Lys Lys  
130 135 140

Glu Lys Arg Ala Thr Ser Trp Pro Asp Leu Ile Ala Lys Val Phe Arg  
145 150 155 160

Ile Asp Val Lys Ala Asp Val Asp Ser Ile His Pro Thr Glu Phe Cys  
165 170 175

His Asn Cys Trp Ser Ile Met His Arg Lys Phe Ser Ser Ala Pro Cys  
180 185 190

Glu Val Tyr Phe Pro Arg Asn Val Thr Met Glu Trp His Pro His Thr  
195 200 205

Pro Ser Cys Asp Ile Cys Asn Thr Ala Arg Arg Gly Leu Lys Arg Lys  
210 215 220

Ser Leu Gln Pro Asn Leu Gln Leu Ser Lys Lys Leu Lys Thr Val Leu  
225 230 235 240

Asp Gln Ala Arg Gln Ala Arg Gln Arg Lys Arg Arg Ala Gln Ala Arg  
245 250 255

Ile Ser Ser Lys Asp Val Met Lys Lys Ile Ala Asn Cys Ser Lys Ile  
260 265 270

His Leu Ser Thr Lys Leu Leu Ala Val Asp Phe Pro Glu His Phe Val  
275 280 285

Lys Ser Ile Ser Cys Gln Ile Cys Glu His Ile Leu Ala Asp Pro Val  
290 295 300

Glu Thr Asn Cys Lys His Val Phe Cys Arg Val Cys Ile Leu Arg Cys  
305 310 315 320

Leu Lys Val Met Gly Ser Tyr Cys Pro Ser Cys Arg Tyr Pro Cys Phe  
325 330 335

Pro Thr Asp Leu Glu Ser Pro Val Lys Ser Phe Leu Ser Val Leu Asn  
340 345 350

Ser Leu Met Val Lys Cys Pro Ala Lys Glu Cys Asn Glu Glu Val Ser  
355 360 365

Leu Glu Lys Tyr Asn His His Ile Ser Ser His Lys Glu Ser Lys Glu  
370 375 380

Ile Phe Val His Ile Asn Lys Gly Gly Arg Pro Arg Gln His Leu Leu  
385 390 395 400

Ser Leu Thr Arg Arg Ala Gln Lys His Arg Leu Arg Glu Leu Lys Leu  
405 410 415

Gln Val Lys Ala Phe Ala Asp Lys Glu Glu Gly Gly Asp Val Lys Ser  
420 425 430

Val Cys Met Thr Leu Phe Leu Leu Ala Leu Arg Ala Arg Asn Glu His  
435 440 445

Arg Gln Ala Asp Glu Leu Glu Ala Ile Met Gln Gly Lys Gly Ser Gly  
450 455 460

Leu Gln Pro Ala Val Cys Leu Ala Ile Arg Val Asn Thr Phe Leu Ser  
465 470 475 480

Cys Ser Gln Tyr His Lys Met Tyr Arg Thr Val Lys Ala Ile Thr Gly  
485 490 495

Arg Gln Ile Phe Gln Pro Leu His Ala Leu Arg Asn Ala Glu Lys Val  
500 505 510

Leu Leu Pro Gly Tyr His His Phe Glu Trp Gln Pro Pro Leu Lys Asn  
515 520 525

Val Ser Ser Ser Thr Asp Val Gly Ile Ile Asp Gly Leu Ser Gly Leu  
530 535 540

Ser Ser Ser Val Asp Asp Tyr Pro Val Asp Thr Ile Ala Lys Arg Phe  
545 550 555 560

Arg Tyr Asp Ser Ala Leu Val Ser Ala Leu Met Asp Met Glu Glu Asp  
565 570 575

Ile Leu Glu Gly Met Arg Ser Gln Asp Leu Asp Asp Tyr Leu Asn Gly  
580 585 590

Pro Phe Thr Val Val Val Lys Glu Ser Cys Asp Gly Met Gly Asp Val  
595 600 605

Ser Glu Lys His Gly Ser Gly Pro Val Val Pro Glu Lys Ala Val Arg  
610 615 620

Phe Ser Phe Thr Ile Met Lys Ile Thr Ile Ala His Ser Ser Gln Asn  
625 630 635 640

Val Lys Val Phe Glu Glu Ala Lys Pro Asn Ser Glu Leu Cys Cys Lys  
645 650 655

Pro Leu Cys Leu Met Leu Ala Asp Glu Ser Asp His Glu Thr Leu Thr  
660 665 670

Ala Ile Leu Ser Pro Leu Ile Ala Glu Arg Glu Ala Met Lys Ser Ser  
675 680 685

Glu Leu Met Leu Glu Leu Gly Gly Ile Leu Arg Thr Phe Lys Phe Ile  
690 695 700

Phe Arg Gly Thr Gly Tyr Asp Glu Lys Leu Val Arg Glu Val Glu Gly  
705 710 715 720

Leu Glu Ala Ser Gly Ser Val Tyr Ile Cys Thr Leu Cys Asp Ala Thr  
725 730 735

Arg Leu Glu Ala Ser Gln Asn Leu Val Phe His Ser Ile Thr Arg Ser  
740 745 750

His Ala Glu Asn Leu Glu Arg Tyr Glu Val Trp Arg Ser Asn Pro Tyr  
755 760 765

His Glu Ser Val Glu Glu Leu Arg Asp Arg Val Lys Gly Val Ser Ala  
770 775 780



Lys Pro Phe Ile Glu Thr Val Pro Ser Ile Asp Ala Leu His Cys Asp  
785 790 795 800

Ile Gly Asn Ala Ala Glu Phe Tyr Lys Ile Phe Gln Leu Glu Ile Gly  
805 810 815

Glu Val Tyr Lys Asn Pro Asn Ala Ser Lys Glu Glu Arg Lys Arg Trp  
820 825 830

Gln Ala Thr Leu Asp Lys His Leu Arg Lys Lys Met Asn Leu Lys Pro  
835 840 845

Ile Met Arg Met Asn Gly Asn Phe Ala Arg Lys Leu Met Thr Lys Glu  
850 855 860

Thr Val Asp Ala Val Cys Glu Leu Ile Pro Ser Glu Glu Arg His Glu  
865 870 875 880

Ala Leu Arg Glu Leu Met Asp Leu Tyr Leu Lys Met Lys Pro Val Trp  
885 890 895

Arg Ser Ser Cys Pro Ala Lys Glu Cys Pro Glu Ser Leu Cys Gln Tyr  
900 905 910

Ser Phe Asn Ser Gln Arg Phe Ala Glu Leu Leu Ser Thr Lys Phe Lys  
915 920 925

Tyr Arg Tyr Glu Gly Lys Ile Thr Asn Tyr Phe His Lys Thr Leu Ala  
930 935 940

His Val Pro Glu Ile Ile Glu Arg Asp Gly Ser Ile Gly Ala Trp Ala  
945 950 955 960

Ser Glu Gly Asn Glu Ser Gly Asn Lys Leu Phe Arg Arg Phe Arg Lys  
965 970 975

Met Asn Ala Arg Gln Ser Lys Cys Tyr Glu Met Glu Asp Val Leu Lys  
980 985 990

His His Trp Leu Tyr Thr Ser Lys Tyr Leu Gln Lys Phe Met Asn Ala  
995 1000 1005

His Asn Ala Leu Lys Thr Ser Gly Phe Thr Met Asn Pro Gln Ala  
1010 1015 1020

Ser Leu Gly Asp Pro Leu Gly Ile Glu Asp Ser Leu Glu Ser Gln  
1025 1030 1035

Asp Ser Met Glu Phe  
1040

<210> 3  
<211> 2414  
<212> DNA  
<213> Homo sapiens

<400> 3  
actctcttta cagtcagcct tctgcttgcc acagtcatag tgggcagtca gtgaatcttc 60  
cccaagtgct gacaattaat acctgggtta gcggcaaaga ttcagagagg cgtgagcagc 120  
ccctctggcc ttcagacaaa aatctacgta ccatcagaaa ctatgtctct gcagatggta 180  
acagtcagta ataacatagc cttaattcag ccaggcttct cactgatgaa ttttgatgga 240  
caagttttct tctttggaca aaaaggctgg cccaaaagat cctgccccac tggagttttc 300  
catctggatg taaagcataa ccatgtcaaa ctgaagccta caattttctc taaggattcc 360  
tgctacctcc ctctctctcg ctaccagcc acttgacat tcaaaggcag cttggagtct 420  
gaaaagcatc aatacatcat ccatggaggg aaaacaccaa acaatgaggt ttcagataag 480  
atztatgtca tgtctattgt ttgcaagaac aacaaaaagg ttacttttcg ctgcacagag 540  
aaagacttgg taggagatgt tcctgaagcc agatatggtc attccattaa tgtggtgtac 600  
agccgagggg aaagtatggg tgctctcttt ggaggacgct catacatgcc ttctaccac 660  
agaaccacag aaaaatggaa tagtgtagct gactgcctgc cctgtgtttt cctggtggat 720  
tttgaatttg ggtgtgtac atcatacatt cttccagaac ttcaggatgg gctatctttt 780  
catgtctcta ttgccaaaaa tgacaccatc tatattttag gaggacattc acttgccaat 840  
aatatccggc ctgccaacct gtacagaata agggttgatc ttcccctggg tagcccagct 900  
gtgaattgca cagtcttgcc aggaggaatc tctgtctcca gtgcaatcct gactcaaact 960  
aacaatgatg aatttggtat tgttggtggc tatcagcttg aaaatcaaaa agaagatgac 1020  
tgcaacatca tctctttaga ggacaacaag atagaaattc gtgagatgga gacccagat 1080  
tggaccccag acattaagca cagcaagata tggtttggaa gcaacacggg aaatggaact 1140

```

gtttttcttg gcataccagg agacaataaa caagttgttt cagaaggatt ctatttctat 1200
atgttgaaat gtgctgaaga tgatactaata gaagagcaga caacattcac aaacagtcaa 1260
acatcaacag aagatccagg ggattccact ccctttgaag actctgaaga attttgtttc 1320
agtgcagaag caaatagttt tgatggtgat gatgaatttg acacctataa tgaagatgat 1380
gaagaagatg agtctgagac aggctactgg attacatgct gccctacttg tgatgtggat 1440
atcaacactt gggtagcatt ctattcaact gagctcaaca aaccgcat gatctactgc 1500
tctcatgggg atgggactg ggtccatgct cagtgcattg atctggcaga acgcacactc 1560
atccatctgt cagcaggaag caacaagtat tactgcaatg agcatgtgga gatagcaaga 1620
gctctacaca ctcccaaag agtcctaccc ttaaaaaagc ctccaatgaa atccctccgt 1680
aaaaaagggt ctggaaaaat cttgactcct gccagaagaat cttttcttag aagggtgttt 1740
gattagtttt gcaaaagcct ttcagattca ggtgtatgga atttttgaat ctatttttaa 1800
aatcataaca ttgattttta aaatacattt ttgtttattt aaaatgccta tgttttcttt 1860
tagttacatg aattaagggc cagaaaaaag tgttttataat gcaatgataa ataaagtcac 1920
tctagaccct atacattttg aaaatatattt acccaaatac tcaatttact aatttattct 1980
tactgagga tttctgatct gattttttat tcaacaaacc ttaaaccacc agaagcagta 2040
ataatcatcg aggtatgttt atatttatta tatgagtctt ggtaacaaat aacctataaa 2100
gtgtttatga caaathtagc caataaagaa attaacaccc aaaagaatta aattgattat 2160
tttggtgcaac ataacaattc ggcagttggc caaaacttaa aagcaagatc tactacatcc 2220
cacattagtg ttctttatat accttcaagc aaccctttgg attatgccca tgaacaagtt 2280
agttttctcat agctttacag atgtagatat aaatataaat atatgtatac atatagatag 2340
ataatgttct cactgacac aaaagaagaa ataaataatc tacatcaaaa aaaaaaaaaa 2400
aaaaaaaaaa aaaa 2414

```

```

<210> 4
<211> 527
<212> PRT
<213> Homo sapiens

```

```

<400> 4

```

```

Met Ser Leu Gln Met Val Thr Val Ser Asn Asn Ile Ala Leu Ile Gln
1           5           10           15

```

Pro Gly Phe Ser Leu Met Asn Phe Asp Gly Gln Val Phe Phe Phe Gly  
20 25 30

Gln Lys Gly Trp Pro Lys Arg Ser Cys Pro Thr Gly Val Phe His Leu  
35 40 45

Asp Val Lys His Asn His Val Lys Leu Lys Pro Thr Ile Phe Ser Lys  
50 55 60

Asp Ser Cys Tyr Leu Pro Pro Leu Arg Tyr Pro Ala Thr Cys Thr Phe  
65 70 75 80

Lys Gly Ser Leu Glu Ser Glu Lys His Gln Tyr Ile Ile His Gly Gly  
85 90 95

Lys Thr Pro Asn Asn Glu Val Ser Asp Lys Ile Tyr Val Met Ser Ile  
100 105 110

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Cys | Lys | Asn | Asn | Lys | Lys | Val | Thr | Phe | Arg | Cys | Thr | Glu | Lys | Asp |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |

Leu Val Gly Asp Val Pro Glu Ala Arg Tyr Gly His Ser Ile Asn Val  
130 135 140

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Tyr | Ser | Arg | Gly | Lys | Ser | Met | Gly | Ala | Leu | Phe | Gly | Gly | Arg | Ser |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |

Tyr Met Pro Ser Thr His Arg Thr Thr Glu Lys Trp Asn Ser Val Ala  
165 170 175

Asp Cys Leu Pro Cys Val Phe Leu Val Asp Phe Glu Phe Gly Cys Ala  
180 185 190

Thr Ser Tyr Ile Leu Pro Glu Leu Gln Asp Gly Leu Ser Phe His Val  
195 200 205

Ser Ile Ala Lys Asn Asp Thr Ile Tyr Ile Leu Gly Gly His Ser Leu  
210 215 220

Ala Asn Asn Ile Arg Pro Ala Asn Leu Tyr Arg Ile Arg Val Asp Leu  
225 230 235 240

Pro Leu Gly Ser Pro Ala Val Asn Cys Thr Val Leu Pro Gly Gly Ile  
245 250 255

Ser Val Ser Ser Ala Ile Leu Thr Gln Thr Asn Asn Asp Glu Phe Val  
260 265 270

Ile Val Gly Gly Tyr Gln Leu Glu Asn Gln Lys Arg Met Ile Cys Asn  
275 280 285

Ile Ile Ser Leu Glu Asp Asn Lys Ile Glu Ile Arg Glu Met Glu Thr  
290 295 300

Pro Asp Trp Thr Pro Asp Ile Lys His Ser Lys Ile Trp Phe Gly Ser  
305 310 315 320

Asn Thr Gly Asn Gly Thr Val Phe Leu Gly Ile Pro Gly Asp Asn Lys  
325 330 335

Gln Val Val Ser Glu Gly Phe Tyr Phe Tyr Met Leu Lys Cys Ala Glu  
340 345 350

Asp Asp Thr Asn Glu Glu Gln Thr Thr Phe Thr Asn Ser Gln Thr Ser  
355 360 365

Thr Glu Asp Pro Gly Asp Ser Thr Pro Phe Glu Asp Ser Glu Glu Phe  
370 375 380

Cys Phe Ser Ala Glu Ala Asn Ser Phe Asp Gly Asp Asp Glu Phe Asp  
385 390 395 400

Thr Tyr Asn Glu Asp Asp Glu Glu Asp Glu Ser Glu Thr Gly Tyr Trp  
405 410 415

Ile Thr Cys Cys Pro Thr Cys Asp Val Asp Ile Asn Thr Trp Val Pro  
420 425 430

Phe Tyr Ser Thr Glu Leu Asn Lys Pro Ala Met Ile Tyr Cys Ser His  
435 440 445

Gly Asp Gly His Trp Val His Ala Gln Cys Met Asp Leu Ala Glu Arg  
450 455 460

Thr Leu Ile His Leu Ser Ala Gly Ser Asn Lys Tyr Tyr Cys Asn Glu  
465 470 475 480

His Val Glu Ile Ala Arg Ala Leu His Thr Pro Gln Arg Val Leu Pro  
485 490 495

Leu Lys Lys Pro Pro Met Lys Ser Leu Arg Lys Lys Gly Ser Gly Lys  
500 505 510

Ile Leu Thr Pro Ala Lys Lys Ser Phe Leu Arg Arg Leu Phe Asp  
515 520 525

<210> 5  
<211> 1816  
<212> DNA  
<213> Homo sapiens

<400> 5  
cttctctgcc agaagatacc atttcaactt taacacagca tgatcgaaac atacaaccaa 60  
acttctcccc gatctgcggc cactggactg cccatcagca tgaaaatttt tatgtattta 120  
cttactgttt ttcttatcac ccagatgatt gggtcagcac tttttgctgt gtatcttcat 180  
agaaggttgg acaagataga agatgaaagg aatcttcatg aagattttgt attcatgaaa 240  
acgatacaga gatgcaacac aggagaaaga tccttatact tactgaactg tgaggagatt 300  
aaaagccagt ttgaaggctt tgtgaaggat ataatgttaa acaaagagga gacgaagaaa 360  
gaaaacagct ttgaaatgca aaaaggatgat cagaatcctc aaattgcggc acatgtcata 420  
agtgaggcca gcagtataaac aacatctgtg ttacagtggg ctgaaaaagg atactacacc 480  
atgagcaaca acttggtaac cctggaaaat gggaaacagc tgaccgttaa aagacaagga 540  
ctctattata tctatgccc agtcaccttc tgttccaatc gggagagctc gagtcaagct 600  
ccatttatag ccagcctctg cctaaagtcc cccggtagat tcgagagaat cttactcaga 660  
gctgcaaata cccacagttc cgccaaacct tgcgggcaac aatccattca cttgggagga 720  
gtatttgaat tgcaaccagg tgcttcgggtg tttgtcaatg tgactgatcc aagccaagtg 780  
agccatggca ctggcttcac gtcctttggc ttactcaaac tctgaacagt gtcaccttgc 840  
aggctgtggt ggagctgacg ctgggagtct tcataatata gcacagcggg taagcccacc 900  
ccctgttaac tgcctattta taaccctagg atcctcctta tggagaacta tttattatac 960  
actccaaggc atgtagaact gtaataagtg aattacaggt cacatgaaac caaaacgggc 1020

```

cctgctccat aagagcttat atatctgaag cagcaacccc actgatgcag acatccagag 1080
agtcctatga aaagacaagg ccattatgca caggttgaat tctgagtaaa cagcagataa 1140
cttgccaagt tcagttttgt ttctttgcgt gcagtgtctt tccatggata atgcatttga 1200
tttatcagtg aagatgcaga agggaaatgg ggagcctcag ctcacattca gttatggttg 1260
actctggggt cctatggcct tgttggaggg ggccaggctc tagaacgtct aacacagtgg 1320
agaaccgaaa cccccccccc ccccccgcc accctctcgg acagttattc attctctttc 1380
aatctctctc tctccatctc tctctttcag tctctctctc tcaacctctt tcttccaatc 1440
tctctttctc aatctctctg tttccctttg tcagtctctt cctccccca gtctctcttc 1500
tcaatcccc tttctaacac acacacacac acacacacac acacacacac acacacacac 1560
acacacacac acacacacac agagtcaggg cgttgctagt cagttctctt ctttccaccc 1620
tgtccctatc tctaccacta tagatgaggg tgaggagtag ggagtgcagc cctgagcctg 1680
cccactctc attacgaaat gactgtattt aaaggaaatc tattgtatct acctgcagtc 1740
tccattgttt ccagagtga a cttgtaatta tcttggttatt tattttttga ataataaaga 1800
cctcttaaca ttaaaa 1816

```

```

<210> 6
<211> 261
<212> PRT
<213> Homo sapiens

```

```

<400> 6

```

```

Met Ile Glu Thr Tyr Asn Gln Thr Ser Pro Arg Ser Ala Ala Thr Gly
1           5           10           15

```

```

Leu Pro Ile Ser Met Lys Ile Phe Met Tyr Leu Leu Thr Val Phe Leu
          20           25           30

```

```

Ile Thr Gln Met Ile Gly Ser Ala Leu Phe Ala Val Tyr Leu His Arg
35           40           45

```

```

Arg Leu Asp Lys Ile Glu Asp Glu Arg Asn Leu His Glu Asp Phe Val
50           55           60

```

```

Phe Met Lys Thr Ile Gln Arg Cys Asn Thr Gly Glu Arg Ser Leu Ser
65           70           75           80

```

Leu Leu Asn Cys Glu Glu Ile Lys Ser Gln Phe Glu Gly Phe Val Lys  
85 90 95

Asp Ile Met Leu Asn Lys Glu Glu Thr Lys Lys Glu Asn Ser Phe Glu  
100 105 110

Met Gln Lys Gly Asp Gln Asn Pro Gln Ile Ala Ala His Val Ile Ser  
115 120 125

Glu Ala Ser Ser Lys Thr Thr Ser Val Leu Gln Trp Ala Glu Lys Gly  
130 135 140

Tyr Tyr Thr Met Ser Asn Asn Leu Val Thr Leu Glu Asn Gly Lys Gln  
145 150 155 160

Leu Thr Val Lys Arg Gln Gly Leu Tyr Tyr Ile Tyr Ala Gln Val Thr  
165 170 175

Phe Cys Ser Asn Arg Glu Ala Ser Ser Gln Ala Pro Phe Ile Ala Ser  
180 185 190

Leu Cys Leu Lys Ser Pro Gly Arg Phe Glu Arg Ile Leu Leu Arg Ala  
195 200 205

Ala Asn Thr His Ser Ser Ala Lys Pro Cys Gly Gln Gln Ser Ile His  
210 215 220

Leu Gly Gly Val Phe Glu Leu Gln Pro Gly Ala Ser Val Phe Val Asn  
225 230 235 240

Val Thr Asp Pro Ser Gln Val Ser His Gly Thr Gly Phe Thr Ser Phe  
245 250 255

Gly Leu Leu Lys Leu  
260

<210> 7  
<211> 24  
<212> PRT  
<213> Artificial sequence

<220>  
<223> Blocking peptide



<400> 7

Ser Ser Lys Thr Thr Ser Val Leu Gln Trp Ala Glu Lys Gly Tyr Tyr  
1 5 10 15

Thr Met Ser Asn Asn Leu Val Thr  
20

<210> 8  
<211> 13  
<212> PRT  
<213> Artificial sequence

<220>  
<223> Blocking peptide

<400> 8

Gln Ile Ala Ala His Val Ile Ser Glu Ala Ser Ser Lys  
1 5 10

<210> 9  
<211> 22  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (21)..(21)  
<223> n is dA

<220>  
<221> misc\_feature  
<222> (22)..(22)  
<223> n is dG

<400> 9  
augucucugc agaugguaac nn

22

<210> 10  
<211> 22  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (21)..(21)  
<223> n is dA

<220>  
<221> misc\_feature  
<222> (22)..(22)  
<223> n is dU

<400> 10  
cuguuaccau cugcagagac nn

22

<210> 11  
<211> 22  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (21)..(21)  
<223> n is dC

<220>  
<221> misc\_feature  
<222> (22)..(22)  
<223> n is dC

<400> 11  
gguaggagau cuuccugaag nn

22

<210> 12  
<211> 24  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> n is dC

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dU

<400> 12  
ggggaugggc acugggucca ugnn

24

<210> 13  
<211> 24  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> n is dC

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dC

<400> 13  
agcauggacc cagugcccau ccnn

24

<210> 14  
<211> 22  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (21)..(21)  
<223> n is dA

<220>  
<221> misc\_feature  
<222> (22)..(22)  
<223> n is dU

<400> 14  
cuguuaccu cugcagagac nn

22

<210> 15  
<211> 24  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> n is dC

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dC

<400> 15  
auggcagccu cuuucccacc cann

24

<210> 16  
<211> 24  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> n is dA

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dU

<400> 16  
gguggguggg aaagaggcug ccnn

24

<210> 17  
<211> 25  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dT

<220>

<221> misc\_feature  
<222> (25)..(25)  
<223> n is dC

<400> 17  
aaacuugcag cucagcaaaa aacnn

25

<210> 18  
<211> 26  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (25)..(25)  
<223> n is dU

<220>  
<221> misc\_feature  
<222> (26)..(26)  
<223> n is dU

<400> 18  
gaguuuuuug cugagcugca aguunn

26

<210> 19  
<211> 26  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (25)..(25)  
<223> n is dU

<220>  
<221> misc\_feature  
<222> (26)..(26)  
<223> n is dU

<400> 19  
gaguuuuuug cugagcugca aguunn

26

<210> 20  
<211> 24

<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> n is dC

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dC

<400> 20  
ucacaaaacc cuggcccaug uunn

24

<210> 21  
<211> 24  
<212> RNA  
<213> Artificial sequence

<220>  
<223> RNA molecule

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> n is dG

<220>  
<221> misc\_feature  
<222> (24)..(24)  
<223> n is dA

<400> 21  
ggaacauggg ccaggguuuu gunn

24